

# TNF & IFN-g 223 genes up-regulated in Y1 cells 16 hr

Fold	Cont	TNF+IFNg	Gene name	Acc#
2.2	50	109	Acid beta glucosidase {IMAGE:331646}	W08086.1
2.1	76	157	AMP deaminase 3 {IMAGE:750480}	AA389897.1
2.1	53	109	Arachidonate 12-lipoxygenase, pseudogene 2 {IMAGE:335369}	W36511.1
2.4	115	281	B-cell leukemia/lymphoma 3 {IMAGE:716976}	AA266002.1
2.1	83	172	B-cell translocation gene 2, anti-proliferative {IMAGE:583186}	AA154848.1
2	53	105	Cathelin-like protein {IMAGE:334755}	W17771.1
2	40	79	CCAAT/enhancer binding protein alpha (C/EBP), related sequence 1 {IM	AA473938.1
2	54	106	Coagulation factor II {IMAGE:350344}	W40725.1
2.2	40	88	Crystallin, gamma S {IMAGE:483498}	AI893668.1
2.2	43	94	Cytochrome P450, 2b13, phenobarbital inducible, type c {IMAGE:316281	AI325330.1
2.1	68	141	EST {IMAGE:474319}	AA038807.1
2.6	409	1083	ESTs {IMAGE:315676}	W09957.1
2.4	434	1047	ESTs {IMAGE:316914}	W11926.1
2.1	51	106	ESTs {IMAGE:317515}	W34080.1
2.1	151	316	ESTs {IMAGE:318969}	W34482.1
2.2	37	82	ESTs {IMAGE:330233}	W14737.1
2.3	66	154	ESTs {IMAGE:334410}	W18278.1
2.7	245	663	ESTs {IMAGE:350182}	W34722.1
2.1	53	111	ESTs {IMAGE:372368}	W53544.1
2	258	517	ESTs {IMAGE:373112}	W62546.1
2	65	133	ESTs {IMAGE:385744}	W64241.1
2.1	65	136	ESTs {IMAGE:403413}	W82905.1
2.9	59	170	ESTs {IMAGE:404589}	W82894.1
2	544	1090	ESTs {IMAGE:419146}	W88005.1
2.1	101	217	ESTs {IMAGE:427480}	AA002452.1
2	324	663	ESTs {IMAGE:436999}	AA002783.1
2.3	69	159	ESTs {IMAGE:459562}	AA027381.1
2.1	61	131	ESTs {IMAGE:466371}	AA033504.1

2.4	121	285	ESTs {IMAGE:467873}	AA036517.1
2	54	109	ESTs {IMAGE:478040}	AI322694.1
2.1	60	124	ESTs {IMAGE:478157}	AA049610.1
2.7	291	772	ESTs {IMAGE:482641}	AA061982.1
2.1	77	158	ESTs {IMAGE:483381}	AI552474.1
2	69	135	ESTs {IMAGE:493212}	AI595201.1
3.2	64	204	ESTs {IMAGE:571219}	AA107718.1
2.2	95	208	ESTs {IMAGE:577555}	AA138025.1
6.2	38	234	ESTs {IMAGE:583566}	AA143986.1
4.6	194	896	ESTs {IMAGE:595951}	AA140542.1
9.7	50	484	ESTs {IMAGE:596863}	AA145136.1
2.5	41	102	ESTs {IMAGE:597493}	AA154419.1
4.9	26	128	ESTs {IMAGE:598824}	AA168416.1
2.1	46	95	ESTs {IMAGE:617402}	AA174434.1
2.1	161	341	ESTs {IMAGE:618611}	AA174941.1
17.1	22	376	ESTs {IMAGE:619723}	AA172624.1
3.6	36	129	ESTs {IMAGE:637888}	AA189578.1
2	50	99	ESTs {IMAGE:640013}	AA198524.1
2.1	82	171	ESTs {IMAGE:669765}	AA230444.1
3.7	266	995	ESTs {IMAGE:678854}	AA250462.1
3.4	209	713	ESTs {IMAGE:751826}	AA396152.1
3.4	41	138	ESTs {IMAGE:761264}	AA388221.1
2.2	53	118	ESTs {IMAGE:775033}	AA387356.1
2.3	68	156	ESTs {IMAGE:775092}	AA386832.1
2	47	96	ESTs {IMAGE:775188}	AA387164.1
2.1	49	102	ESTs {IMAGE:775208}	AA387199.1
2	45	88	ESTs {IMAGE:792921}	AA415579.1
2	61	122	ESTs {IMAGE:793692}	AA415691.1
2	53	107	ESTs {IMAGE:804793}	AA472833.1
2.3	104	238	ESTs {IMAGE:805129}	AA473234.1
3.4	45	153	ESTs {IMAGE:805386}	AA473205.1
4	393	1563	ESTs {IMAGE:817924}	AA466508.1

2.4	61	146	ESTs {IMAGE:818790}	AA467382.1
2	38	75	ESTs {IMAGE:831741}	AA444941.1
2.3	38	87	ESTs {IMAGE:833477}	AA465972.1
2.6	62	159	ESTs {IMAGE:834617}	AA467546.1
2.1	48	100	ESTs {IMAGE:835536}	AA422515.1
2.1	43	91	ESTs {IMAGE:836133}	AA426897.1
3.5	59	209	ESTs {IMAGE:837565}	AA432818.1
2	47	95	ESTs {IMAGE:845839}	AA416091.1
2	80	158	ESTs {IMAGE:872986}	AA466235.1
2.1	99	207	ESTs {IMAGE:876106}	AA475528.1
2	50	98	ESTs {IMAGE:904836}	AA522117.1
3.6	118	426	ESTs {IMAGE:945014}	AA537234.1
2.8	48	132	ESTs {IMAGE:961094}	AA547202.1
2.1	147	310	ESTs, Highly similar to CYSTATHIONINE GAMMA-LYASE [Homo sapiens]	AI892342.1
2.5	59	148	ESTs, Highly similar to CYSTATHIONINE GAMMA-LYASE [Homo sapiens]	AA245993.1
5.7	55	313	ESTs, Highly similar to HYPOTHETICAL 103.5 KD PROTEIN IN RAD26-C	AI639807.1
4	580	2318	ESTs, Highly similar to INTERFERON-INDUCIBLE PROTEIN [Rattus norvegicus]	AA048837.1
6.3	40	253	ESTs, Highly similar to LYSYL-TRNA SYNTHETASE [Cricetulus longicaudatus]	AA175815.1
4.1	165	683	ESTs, Highly similar to NUCLEAR FACTOR NF-KAPPA-B P100 SUBUNIT	AI385509.1
3	149	449	ESTs, Highly similar to SERINE HYDROXYMETHYLTRANSFERASE, MITOCHONDRIAL	AA208877.1
2.2	152	336	ESTs, Highly similar to UTP--GLUCOSE-1-PHOSPHATE URIDYLTRANSFERASE	AA473123.1
2.3	389	888	ESTs, Highly similar to 54 kDa oligoadenylate synthetase-like protein p1	AA123007.1
2.3	49	111	ESTs, Highly similar to FLASH [Mus musculus] {IMAGE:833829}	AA412925.1
2.8	218	603	ESTs, Highly similar to KIAA0650 protein [H.sapiens] {IMAGE:621629}	AA183687.1
3.2	62	197	ESTs, Highly similar to major vault protein [R.norvegicus] {IMAGE:6395}	AA200827.1
2	67	131	ESTs, Highly similar to molybdopterin-synthase large subunit [Mus musculus]	AI450527.1
2.2	45	100	ESTs, Highly similar to myomegalin [R.norvegicus] {IMAGE:315787}	W12392.1
2.2	50	110	ESTs, Highly similar to NY-REN-2 antigen [H.sapiens] {IMAGE:945689}	AA546195.1
2	47	92	ESTs, Highly similar to pirin [H.sapiens] {IMAGE:332396}	W08720.1
2.6	142	373	ESTs, Highly similar to probable nocturnin protein [Mus musculus] {IMAGE:1133.1}	AA051133.1
2.6	71	185	ESTs, Highly similar to putative transcription factor [H.sapiens] {IMAGE:436.1}	AA467436.1
2	208	410	ESTs, Highly similar to unknown [H.sapiens] {IMAGE:336564}	AI390954.1

2.3	132	298	ESTs, Highly similar to unknown [H.sapiens] {IMAGE:492447}	AA097194.1
3.2	87	277	ESTs, Moderately similar to ACTIN-LIKE PROTEIN 14D [Drosophila mel	AA451385.1
2	46	92	ESTs, Moderately similar to GEPHYRIN [Rattus norvegicus] {IMAGE:61	AI893727.1
2	49	100	ESTs, Moderately similar to C11D2.4 [C.elegans] {IMAGE:441807}	AA014053.1
2	50	101	ESTs, Moderately similar to homocysteine respondent protein HCYP2 [I	W20810.1
2	337	665	ESTs, Moderately similar to KIAA0651 protein [H.sapiens] {IMAGE:6952	AA220487.1
2.1	68	140	ESTs, Moderately similar to probable ligand-binding protein RYD5 [R.n	AA412821.1
2.1	61	130	ESTs, Moderately similar to SARCOLIPIN [H.sapiens] {IMAGE:318692}	W11587.1
2.9	61	174	ESTs, Moderately similar to unknown [H.sapiens] {IMAGE:761038}	AA387579.1
2.1	59	124	ESTs, Weakly similar to CYP2C39 [M.musculus] {IMAGE:367514}	AI427394.1
2.3	39	90	ESTs, Weakly similar to cytohesin 2 [M.musculus] {IMAGE:779020}	AA413956.1
2.1	85	177	ESTs, Weakly similar to CYTOSKELETAL P17 PROTEIN [Dictyostelium	AA017875.1
2.2	59	131	ESTs, Weakly similar to DUAL SPECIFICITY PROTEIN PHOSPHATASE 1	AA119029.1
2.2	623	1381	ESTs, Weakly similar to GARP PROTEIN PRECURSOR [H.sapiens] {IMA	AA048874.1
2.3	43	97	ESTs, Weakly similar to gob-5 [M.musculus] {IMAGE:351090}	W41083.1
2.1	71	150	ESTs, Weakly similar to HYPOTHETICAL 11.4 KD PROTEIN C13G6.04 IN	W11535.1
2.2	45	98	ESTs, Weakly similar to keratin 10, type I, cytoskeletal [H.sapiens] {IMA	W18457.1
3.1	310	950	ESTs, Weakly similar to LYMPHOCYTE ANTIGEN LY-6A.2/LY-6E.1 PREC	AA472994.1
3.5	81	283	ESTs, Weakly similar to lysophospholipase I [M.musculus] {IMAGE:596	AA138854.1
2.5	705	1750	ESTs, Weakly similar to mannose-binding lectin associated serine protei	AA209006.1
2.3	107	249	ESTs, Weakly similar to mCAC [M.musculus] {IMAGE:350881}	W40994.1
2.6	81	212	ESTs, Weakly similar to ORF YJL004c [S.cerevisiae] {IMAGE:933229}	AA541946.1
2	57	113	ESTs, Weakly similar to Pro-Pol-dUTPase polypeptide [M.musculus] {IM	AA518000.1
2.1	180	371	ESTs, Weakly similar to SDP8 [M.musculus] {IMAGE:747180}	AA260397.1
2.2	66	143	ESTs, Weakly similar to similar to alpha/beta hydrolase fold [C.elegans]	W09936.1
2.1	76	161	ESTs, Weakly similar to similar to leucyl-tRNA synthetase [C.elegans] {I	AA397172.1
2.3	566	1300	ESTs, Weakly similar to tazarotene-induced gene 2 [H.sapiens] {IMAGE	AA220505.1
2.3	315	711	ESTs, Weakly similar to Trif [M.musculus] {IMAGE:523380}	AA080169.1
2.4	298	724	ESTs, Weakly similar to UBIQUITIN-CONJUGATING ENZYME E2-16 KD	AA050167.1
2.5	591	1467	Extracellular matrix protein 1 {IMAGE:874833}	AA474897.1
2.2	428	926	Glutamate oxaloacetate transaminase 1, soluble {IMAGE:481381}	AA060494.1

2.2	45	97	Hemochromatosis {IMAGE:313224}	W11889.1
12.5	37	464	Histocompatibility 2, class II antigen A, alpha {IMAGE:747378}	AA272807.1
3.1	62	190	Histocompatibility 2, class II antigen A, beta 1 {IMAGE:618271}	AA175329.1
3.9	59	231	Histocompatibility 2, class II, locus Ma {IMAGE:669933}	AA230649.1
2	329	660	Histocompatibility 2, K region {IMAGE:330336}	W14540.1
2.2	143	314	Histocompatibility 2, Q region locus 7 {IMAGE:596447}	AA122791.1
2.2	311	688	Histocompatibility 2, Q region locus 7 {IMAGE:637849}	AA186012.1
2	560	1133	Histocompatibility 2, T region locus 22 {IMAGE:920425}	AA538511.1
2.1	107	225	Homolog of human MLLT2 unidentified gene {IMAGE:750930}	AA289139.1
6.7	36	240	Interferon activated gene 203 {IMAGE:620541}	AA174447.1
4.1	273	1123	Interferon regulatory factor 1 {IMAGE:777393}	AA276697.1
3.4	801	2763	Lymphocyte antigen 6 complex {IMAGE:580715}	AA145865.1
2.7	539	1462	Lymphocyte antigen 6 complex, locus C {IMAGE:425855}	AA000712.1
5.8	37	216	M.musculus mRNA for caspase-11 {IMAGE:618272}	AA175651.1
2.5	534	1329	M.musculus mRNA for mTGIF protein {IMAGE:474339}	AA038657.1
3.3	322	1066	M.musculus mRNA for mTGIF protein {IMAGE:722623}	AA260654.1
2.3	61	140	M.musculus spasmodic polypeptide (SP) mRNA {IMAGE:314078}	W09829.1
2.1.	47	99	Metallothionein 2 {IMAGE:427339}	AA003303.1
49.5	62	3070	Monokine induced by gamma interferon {IMAGE:874758}	AA472492.1
3	557	1655	Mouse chromatin nonhistone high mobility group protein (HGM-I(Y), co	AA538243.1
2.3	94	216	Mus musculus A10 mRNA, partial cds {IMAGE:333376}	W15888.1
2.2	401	902	Mus musculus A10 mRNA, partial cds {IMAGE:385441}	W61383.1
2	72	145	Mus musculus Ankhzn mRNA, complete cds {IMAGE:894596}	AA517717.1
2	349	710	Mus musculus asparagine synthetase mRNA, complete cds {IMAGE:33:	W29492.1

2	147	287	Mus musculus BS4 peptide mRNA, complete cds {IMAGE:459091}	AA024028.1
2	162	317	Mus musculus CIPER protein (Ciper) mRNA, complete cds {IMAGE:355}	W47752.1
2.1	64	137	Mus musculus classII MHC transactivator CIITA mRNA, complete cds {I	AA178010.1
2.1	400	836	Mus musculus G-protein-like LRG-47 mRNA, complete cds {IMAGE:651}	AA214784.1
2.1	36	74	Mus musculus IER5 (Ier5) mRNA, complete cds {IMAGE:643745}	AA203798.1
2.4	57	136	Mus musculus interferon regulatory factor 5 (mirf5) mRNA, complete cd	AA185025.1
53.7	29	1558	Mus musculus interferon-induced guanylate binding protein GBP-2 (Gb	AA153021.1
2.4	44	107	Mus musculus intersectin-EH binding protein Ibp2 mRNA, partial cds {I	AA413761.1
2.1	141	291	Mus musculus ion channel homolog RIC mRNA, complete cds {IMAGE:	W98807.1
2.2	59	127	Mus musculus major histocompatibility complex region NG27, NG28, R	W13905.1
2.5	36	90	Mus musculus mRNA for DNA helicase Q1, complete cds {IMAGE:3304}	AI390823.1
6.4	39	249	Mus musculus mRNA for GTP1 protein {IMAGE:621346}	AA178613.1
20.2	35	708	Mus musculus mRNA for ILGP protein {IMAGE:747208}	AA260490.1
20.9	50	1044	Mus musculus mRNA for ILGP protein {IMAGE:764385}	AA277451.1
2	42	82	Mus musculus mRNA for MD-1, complete cds {IMAGE:402964}	W80147.1
2	49	99	Mus musculus peroxisomal biogenesis factor (Pex11a) mRNA, complet	AA237383.1
10	81	809	Mus musculus putative purine nucleotide binding protein mRNA, compl	AA240404.1
2.3	128	288	Mus musculus secreted carbonic anhydrase isozyme VI precursor, mRi	AI327498.1
2.5	535	1362	Myelocytomatosis oncogene {IMAGE:441346}	AA009268.1
2.7	234	625	Myeloid differentiation primary response gene 118 {IMAGE:386412}	W64388.1
3.2	520	1650	MYELOID DIFFERENTIATION PRIMARY RESPONSE PROTEIN MYD116 {	AA050417.1
2	352	719	NAD-DEPENDENT METHYLENETETRAHYDROFOLATE DEHYDROGENA	W84014.1
3.2	507	1629	Protease (prosome, macropain) 28 subunit, alpha {IMAGE:678768}	AA239485.1
2	83	162	Protein tyrosine phosphatase, non-receptor type 2 {IMAGE:437517}	AA003599.1
2.2	48	105	Public domain EST {IMAGE:314509}	W10776.1
2.2	49	110	Public domain EST {IMAGE:314524}	W10780.1
2.3	53	122	Public domain EST {IMAGE:330298}	W13940.1
2	53	106	Public domain EST {IMAGE:350327}	W40724.1
2.4	82	194	Public domain EST {IMAGE:353202}	W41810.1

2.1	53	109	Public domain	EST {IMAGE:367869}	W54520.1
2.1	42	88	Public domain	EST {IMAGE:372366}	W53543.1
2	63	128	Public domain	EST {IMAGE:374745}	W59461.1
2.1	39	83	Public domain	EST {IMAGE:390150}	W70677.1
2.2	41	89	Public domain	EST {IMAGE:403961}	W82923.1
2	131	256	Public domain	EST {IMAGE:439362}	AI466638.1
2.1	47	101	Public domain	EST {IMAGE:442201}	AA013653.1
2.2	58	129	Public domain	EST {IMAGE:456831}	AA023545.1
2	57	114	Public domain	EST {IMAGE:459599}	AA027528.1
2	96	196	Public domain	EST {IMAGE:463183}	AA030360.1
2	50	101	Public domain	EST {IMAGE:465170}	AA030746.1
2.3	48	109	Public domain	EST {IMAGE:466376}	AA032359.1
2	44	90	Public domain	EST {IMAGE:478309}	AA049436.1
2	51	101	Public domain	EST {IMAGE:479480}	AA049260.1
4.3	58	252	Public domain	EST {IMAGE:482196}	AA059908.1
2.3	78	176	Public domain	EST {IMAGE:482817}	AA062120.1
3.3	93	303	Public domain	EST {IMAGE:574439}	AA118291.1
4.7	108	507	Public domain	EST {IMAGE:596231}	AA122443.1
13.8	27	372	Public domain	EST {IMAGE:597020}	AA138757.1
4.4	121	536	Public domain	EST {IMAGE:617529}	AA174721.1
4.6	77	355	Public domain	EST {IMAGE:618199}	AA175243.1
2.1	66	138	Public domain	EST {IMAGE:619596}	AA175803.1
2	82	162	Public domain	EST {IMAGE:620030}	AA178100.1
3.5	93	324	Public domain	EST {IMAGE:621742}	AA177584.1
6.3	27	169	Public domain	EST {IMAGE:636640}	AA204588.1
2.2	41	89	Public domain	EST {IMAGE:652811}	AA217625.1
4	35	141	Public domain	EST {IMAGE:653161}	AI390695.1
3.1	169	521	Public domain	EST {IMAGE:733734}	AA272876.1
2.1	93	193	Public domain	EST {IMAGE:760824}	AA386627.1
2.8	47	130	Public domain	EST {IMAGE:760917}	AA386815.1
2.8	57	159	Public domain	EST {IMAGE:761009}	AA387552.1
2.4	62	147	Public domain	EST {IMAGE:774996}	AA387314.1

4.6	50	231	Public domain EST {IMAGE:775154}	AI604749.1
2.2	44	97	Public domain EST {IMAGE:778923}	AA414282.1
2	44	86	Public domain EST {IMAGE:779453}	AA414212.1
2.2	67	146	Public domain EST {IMAGE:805306}	AA473329.1
2.5	71	180	Public domain EST {IMAGE:948845}	AA544859.1
2.7	86	229	Recombining binding protein suppressor of hairless (Drosophila) {IMA	W97158.1
2.3	39	88	SAPK/Erk/kinase 1 {IMAGE:778459}	AA413508.1
7.4	116	859	Signal transducer and activator of transcription 1 {IMAGE:749544}	AA388678.1
2.2	52	113	Thymus cell antigen 1, theta {IMAGE:316409}	W13151.1
15.7	40	629	Transporter 1, ABC (ATP binding cassette) {IMAGE:576775}	AA123837.1
2.3	45	104	Transthyretin {IMAGE:335319}	W17647.1
7.7	198	1527	TRYPTOPHANYL-TRNA SYNTHETASE {IMAGE:367765}	W53959.1
2.6	401	1028	Tumor necrosis factor Induced protein 2 {IMAGE:949472}	AA537672.1
2.2	80	175	Zinc finger protein 35 {IMAGE:422121}	AI893996.1
2.3	82	190	Zinc transporter 4 {IMAGE:643619}	AA203866.1



Tg 1hr 72 genes up >= 2-fold 022PAV3Q (Exp M subm 2/17/00)

Fold	Cont	Tg	Gene name	Acc#
2.7	527	1434	apoptosis Inhibitor 1 {IMAGE:640046}	AA197349.1
2.1	279	581	B-cell leukemia/lymphoma 10 {IMAGE:355209}	W47752.1
2.5	2462	6113	beta-2 microglobulin {IMAGE:572542}	AA109951.1
2.1	609	1263	capping protein alpha 1 {IMAGE:779754}	AA414612.1
3.1	125	388	caspase 11 {IMAGE:618272}	AA175651.1
2.1	1054	2215	cell division cycle 42 homolog (S. cerevisiae) {IMAGE:721518}	AA266975.1
3.4	321	1087	ESTs {IMAGE:523814}	AA067146.1
3.4	190	639	ESTs {IMAGE:583566}	AA143986.1
9.6	369	3541	ESTs {IMAGE:595951}	AA140542.1
2	150	298	ESTs {IMAGE:596556}	AA146363.1
3.9	165	647	ESTs {IMAGE:596863}	AA145136.1
2.5	108	265	ESTs {IMAGE:597031}	AA145466.1
5.6	148	827	ESTs {IMAGE:619723}	AA172624.1
3.5	125	441	ESTs {IMAGE:636640}	AA204588.1
2	556	1133	ESTs {IMAGE:636827}	AA183302.1
5.7	366	2072	ESTs {IMAGE:678854}	AA250462.1
3.1	306	954	ESTs {IMAGE:751826}	AA396152.1
3.3	162	542	ESTs {IMAGE:777379}	AA277040.1
2.5	368	928	ESTs {IMAGE:876257}	AA476183.1
2	237	484	ESTs, Highly similar to HYPOTHETICAL 103.5 KD PROTEIN IN RAD26-GI	AI639807.1
2.2	224	502	ESTs, Highly similar to INTERFERON-ACTIVABLE PROTEIN 204 [Mus	AI591873.1
2.8	866	2458	ESTs, Highly similar to INTERFERON-INDUCIBLE PROTEIN [Rattus norv	AA048837.1
2.1	243	511	ESTs, Highly similar to AE-1 binding protein AEBP2 [M.musculus] {IMAG	AA416308.1
2.2	556	1207	ESTs, Highly similar to Bclight protein [R.norvegicus] {IMAGE:622724}	AA177555.1
2.5	426	1070	ESTs, Highly similar to probable nocturnin protein [M.musculus] {IMAGE	AA051133.1
7.2	293	2099	ESTs, Highly similar to purine nucleotide binding protein [M.musculus] {I	AA240404.1
2.6	423	1108	ESTs, Highly similar to spliceosomal protein SAP 155 [H.sapiens] {IMAGI	AA200809.1

3.6	124	451	ESTs, Moderately similar to ACTIN-LIKE PROTEIN 14D [Drosophila mela	AA451385.1
3.3	663	2170	ESTs, Moderately similar to CHLORINE CHANNEL PROTEIN P64 [Bos ta	AA466508.1
2	197	391	ESTs, Moderately similar to CHLORINE CHANNEL PROTEIN P64 [Bos ta	AA537234.1
2.4	188	442	ESTs, Moderately similar to PLASMA-CELL MEMBRANE GLYCOPROTEI	AA016397.1
2.2	680	1508	ESTs, Moderately similar to PRE-MRNA SPLICING FACTOR SRP20 [Hom	W83358.1
2.2	330	740	ESTs, Weakly similar to CCAAT-BINDING TRANSCRIPTION FACTOR SUE	AA413894.1
2.9	785	2288	ESTs, Weakly similar to LYMPHOCYTE ANTIGEN LY-6A.2/LY-6E.1 PRECI	AA472994.1
2.5	160	403	ESTs, Weakly similar to lysophospholipase I [M.musculus] {IMAGE:5964	AA153793.1
2.8	1172	3251	ESTs, Weakly similar to mannose-binding lectin associated serine protea	AA209006.1
2.8	625	1738	ESTs, Weakly similar to SDP8 [M.musculus] {IMAGE:747180}	AA260397.1
2.1	413	871	ESTs, Weakly similar to Trif [M.musculus] {IMAGE:523380}	AA080169.1
3.3	214	706	expressed sequence tag mouse EST 25 {IMAGE:638284}	AA200747.1
2.1	713	1511	interferon activated gene 203 {IMAGE:618572}	AA170251.1
2.7	255	697	interferon activated gene 203 {IMAGE:620541}	AA174447.1
4.8	1249	5945	interferon inducible protein 1 {IMAGE:651927}	AA214784.1
9.6	411	3937	interferon regulatory factor 1 {IMAGE:777393}	AA276697.1
2.8	2057	5726	lymphocyte antigen 6 complex {IMAGE:580715}	AA145865.1
2.5	792	1977	lymphocyte antigen 6 complex, locus C {IMAGE:425855}	AA000712.1
4.4	225	980	monokine induced by gamma interferon {IMAGE:874758}	AA472492.1
21	204	4277	Mus musculus Interferon-induced guanylate binding protein GBP-2 (Gbp	AA153021.1
2.8	279	777	Mus musculus mRNA for GTPi protein {IMAGE:621346}	AA178613.1
2	1359	2725	Mus musculus mRNA for heterogeneous nuclear ribonucleoprotein H {IM	AA003085.1
4.4	125	549	Mus musculus mRNA for IIGP protein {IMAGE:747208}	AA260490.1
5.6	229	1293	Mus musculus mRNA for IIGP protein {IMAGE:764385}	AA277451.1
2	268	546	myeloid differentiation primary response gene 88 {IMAGE:572317}	AA110412.1
4.8	228	1096	nuclear factor of kappa light chain gene enhancer in B-cells Inhibitor, alp	AA204405.1
2	606	1236	protein kinase C, delta {IMAGE:777583}	AA276844.1
3.6	184	664	Public domain EST {IMAGE:482196}	AA059908.1
2.9	132	379	Public domain EST {IMAGE:596231}	AA122443.1

5.8	105	614	Public domain EST {IMAGE:597020}	AA138757.1
7.1	221	1571	Public domain EST {IMAGE:617529}	AA174721.1
2	122	247	Public domain EST {IMAGE:619686}	AA175973.1
3.4	138	471	Public domain EST {IMAGE:653161}	AI390695.1
2	215	425	Public domain EST {IMAGE:669831}	AA253750.1
2.3	411	935	Public domain EST {IMAGE:733734}	AA272876.1
2.1	383	801	Public domain EST {IMAGE:850178}	AA451403.1
2.7	1687	4542	Public domain EST {IMAGE:874030}	AA472200.1
2.6	579	1531	purine-nucleoside phosphorylase {IMAGE:607469}	AA172867.1
7.8	222	1740	signal transducer and activator of transcription 1 {IMAGE:749544}	AA388678.1
3	617	1833	signal transducer and activator of transcription 3 {IMAGE:440571}	AA014643.1
3	666	1990	small inducible cytokine subfamily D, 1 {IMAGE:761708}	AA272232.1
4.4	138	613	transporter 1, ABC (ATP binding cassette) {IMAGE:576775}	AA123837.1
2.2	329	719	tryptophanyl-tRNA synthetase {IMAGE:367765}	W53959.1
2	357	717	Y box protein 3 {IMAGE:421546}	W97158.1
2.4	944	2275	zinc finger protein 36 {IMAGE:641523}	AA200282.1